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PubMed Nucleotide Protein Genome Structure PopSet Taxonomy OMIM Boo

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☐ 1: Q04791. FATTY ACYL-COA HY...[gi:730714]

NEW Links

LOCUS SASB_ANAPL 557 aa linear VRT 30-MAY-2000
DEFINITION FATTY ACYL-COA HYDROLASE PRECURSOR, MEDIUM CHAIN (THIOESTERASE B).
ACCESSION Q04791
VERSION Q04791 GI:730714
DBSOURCE swissprot: locus SASB_ANAPL, accession Q04791;
class: standard.
created: Feb 1, 1995.
sequence updated: Feb 1, 1995.
annotation updated: May 30, 2000.
xrefs: gi: 213100, gi: 213101
xrefs (non-sequence databases): HSSP P21836, PFAM PF00135, PROSITE
PS00122, PROSITE PS00941
KEYWORDS Fatty acid biosynthesis; Hydrolase; Signal.
SOURCE Anas platyrhynchos.
ORGANISM Anas platyrhynchos
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Anseriformes; Anatidae; Anas.
REFERENCE 1 (residues 1 to 557)
AUTHORS Hwang, C.S. and Kolattukudy, P.E.
TITLE Molecular cloning and sequencing of thioesterase B cDNA and
stimulation of expression of the thioesterase B gene associated
with hormonal induction of peroxisome proliferation
J. Biol. Chem. 268 (19), 14278-14284 (1993)
JOURNAL 93300823
MEDLINE 8314791
PUBMED
REMARK SEQUENCE FROM N.A., AND SEQUENCE OF 26-65.
TISSUE=UROPYGIAL GLAND
COMMENT

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collaboration between the Swiss Institute of Bioinformatics and
the EMBL outstation - the European Bioinformatics Institute.
The original entry is available from <http://www.expasy.ch/sprot>
and <http://www.ebi.ac.uk/sprot>

[FUNCTION] FATTY ACID BIOSYNTHESIS CHAIN TERMINATION AND RELEASE OF
THE FREE FATTY ACID PRODUCT IS ACHIEVED BY HYDROLYSIS OF THE THIO
ESTER BY A THIOESTERASE. THIS THIOESTERASE MAY BE ASSOCIATED WITH
PEROXISOME PROLIFERATION AND MAY PLAY A ROLE IN THE PRODUCTION OF
3-HYDROXY FATTY ACID DIESTER PHEROMONES.
[CATALYTIC ACTIVITY] OLEOYL-[ACYL-CARRIER PROTEIN] + H(2)O =
ACYL-CARRIER PROTEIN + OLEATE.
[TISSUE SPECIFICITY] HIGHEST LEVELS IN UROPYGIAL GLAND, MUCH LOWER
IN LIVER AND KIDNEY.
[SIMILARITY] BELONGS TO THE TYPE-B CARBOXYLESTERASE/LIPASE FAMILY.
FEATURES
source Location/Qualifiers
1..557
/organism="Anas platyrhynchos"
/db_xref="taxon:8839"

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FEATURES

source

Location/Qualifiers

1..557

/organism="Anas platyrhynchos"

/db_xref="taxon:8839"

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Region 26..557
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Site 345
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Site 460
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Site 476
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 ORIGIN

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Revised: July 5, 2002.

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Aug 28 2002 15:52:55